•	changed a file from non-ASCII to ASCII Changed a file from non-ASCII to ASCII
	Changed the margins in cases where the sequence text was "wrapped" down to the next lin
	Edited a format error in the Current Application Data section, specifically:
_	Edited a format entir in the Current Application Data Socion, Specifically.
	Edited the Current Application Data section with the actual current number. The number inputted by the prior application data; or other
A	Added the mandatory heading and subheadings for "Current Application Data".
Ε	Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integ
C	Changed the spelling of a mandatory field (the headings or subheadings), specifically:
C	Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
lr	nserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
C	corrected subheading placement. All responses must be on the same line as each subheading. If the pplicant placed a response below the subheading, this was moved to its appropriate place.
li	nserted colons after headings/subheadings. Headings edited included:
_	nserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically:
_ _ _	Deleted extra, invalid, headings used by an applicant, specifically:
- - - !	Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end
	Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of page numbers throughout text; other invalid text, such as
	Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end in page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically:
	Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically:
	Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically:
	Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end page numbers throughout text; other invalid text, such as Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically:

^{*}Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



PCT09

RAW SEQUENCE LISTING DATE: 03/07/2002 PATENT APPLICATION: US/09/913,756 TIME: 18:40:43

Input Set : A:\PTO.AMC.txt

```
3 <110> APPLICANT: Chiari, Rita
          Coulie, Pierre
  5
          Boon-Falleur, Thierry
  7 <120> TITLE OF INVENTION: TYROSINE KINASE RECEPTOR EPHA3 ANTIGENIC PEPTIDES
  9 <130> FILE REFERENCE: L0461/7121
 11 <140> CURRENT APPLICATION NUMBER: US 09/913,756
 12 <141> CURRENT FILING DATE: 2000-02-18
 14 <150> PRIOR APPLICATION NUMBER: US 60/121,170
 15 <151> PRIOR FILING DATE: 1999-02-22
 17 <150> PRIOR APPLICATION NUMBER: US 60/158,566
 18 <151> PRIOR FILING DATE: 1999-10-08
 20 <160> NUMBER OF SEQ ID NOS: 63
 22 <170> SOFTWARE: FastSEQ for Windows Version 3.0
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 33 <223> OTHER INFORMATION: synthetic oligo(dT) primer
 35 <220> FEATURE:
 36 <221> NAME/KEY: unsure
 37 <222> LOCATION: 43..43
 38 <223> OTHER INFORMATION: n = a, c, g or t
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> 41 ataagaatgc ggccgctaaa ctatttttt tttttttt tvn^{\prime}
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 45 <211> LENGTH: 3149
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 49 <220> FEATURE:
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 56
                                                 Met Asp Cys Gln Leu
 57
 59 tee ate ete ett ete age tge tet gtt ete gae age tte ggg gaa
                                                                           163
 60 Ser Ile Leu Leu Leu Ser Cys Ser Val Leu Asp Ser Phe Gly Glu
 61
                      10
                                          15
 63 ctg att ccg cag cct tcc aat gaa gtc aat cta ctg gat tca aaa aca
                                                                           211
```

Input Set : A:\PTO.AMC.txt

64 65	Leu	Ile	Pro	Gln 25	Pro	Ser	Asn	Glu	Val 30	Asn	Leu	Leu	Asp	Ser 35	Lys	Thr	
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68					-								His			_	233
	TIE	GIII		GIU	пец	GLY	пр	45	261	тут	PIU	261	50	GLY	тъ	Giu	
69			40											4_			
71			_			_	_						agg			_	307
72	Glu	Ile	Ser	Gly	Val	Asp	Glu	His	Tyr	Thr	Pro	Ile	Arg	Thr	${ t Tyr}$	Gln	
73		55					60					65					
75	gtg	tgc	aat	gtc	atg	gac	cac	agt	caa	aac	aat	tgg	ctg	aga	aca	aac	355
76	Val	Cys	Asn	Val	Met	Asp	His	Ser	Gln	Asn	Asn	Trp	Leu	Arg	\mathtt{Thr}	Asn	
77	70					75					80					85	
79	tgg	gtc	ccc	agg	aac	tca	gct	cag	aag	att	tat	gtg	gag	ctc	aaq	ttc	403
80													Ğlu				
81	_			_	90				-	95		•			100		
83	act	cta	cga	gac		aat	age	att	cca		att	tta	gga	act		ааσ	451
84			_	_	_		-			_	_		Gly		-	-	431
	1111	Бец	ALY	105	Cys	ASII	361	116		пеа	vai	пец	GLy	1115	Cys	шуз	
85				-					110								400
87					-			-			-	_	gat				499
88	GIu	Thr		Asn	Leu	Tyr	Tyr		GIu	Ser	Asp	Asp	Asp	His	GTA	Val	
89			120					125					130				
91			_			_			_		-		att	_	-	-	547
92	Lys	Phe	Arg	Glu	His	Gln	Phe	Thr	Lys	Ile	Asp	Thr	Ile	Ala	Ala	Asp	
93		135					140					145					
95	gaa	agt	ttc	act	caa	atg	gat	ctt	ggg	gac	cgt	att	ctg	aag	ctc	aac	595
95 96													ctg Leu				595
																	595
96	Glu 150	Ser	Phe	Thr	Gln	Met 155	Asp	Leu	Gly	Asp	Arg 160	Ile	Leu	Lys	Leu	Asn 165	
96 97 99	Glu 150 act	Ser gag	Phe att	Thr aga	Gln gaa	Met 155 gta	Asp	Leu cct	Gly gtc	Asp	Arg 160 aag	Ile aag	Leu gga	Lys	Leu tat	Asn 165 ttg	595 643
96 97 99 100	Glu 150 act	Ser gag	Phe att	Thr aga	Gln gaa Glu	Met 155 gta Val	Asp	Leu cct	Gly gtc	Asp aac Asn	Arg 160 aag Lys	Ile aag	Leu gga	Lys	Leu tat Tyr	Asn 165 ttg Leu	
96 97 99 100 101	Glu 150 act Thr	Ser gag Glu	Phe att ı Ile	Thr aga Arg	Gln gaa Glu 170	Met 155 gta Val	Asp ggt Gly	Leu cct Pro	Gly gtc Val	Asp aac Asn 175	Arg 160 aag Lys	Ile aag Lys	Leu gga Gly	Lys ttt Phe	Leu tat Tyr 180	Asn 165 ttg Leu	643
96 97 99 100 101 103	Glu 150 act Thr	Ser gag Glu	Phe att Ile	Thr aga Arg	Gln gaa Glu 170 gtt	Met 155 gta Val	Asp ggt Gly gct	Leu cct Pro	Gly gtc Val gtt	Asp aac Asn 175	Arg 160 aag Lys	Ile aag Lys	Leu gga Gly tct	Lys ttt Phe	tat Tyr 180	Asn 165 ttg Leu	
96 97 99 100 101 103 104	Glu 150 act Thr	Ser gag Glu	Phe att Ile	Thr aga Arg agat Asp	gaa Glu 170 gtt Val	Met 155 gta Val	Asp ggt Gly gct	Leu cct Pro	gtc ytal yal gtt Val	Asp aac Asn 175 gcc Ala	Arg 160 aag Lys	Ile aag Lys	Leu gga Gly tct	ttt Phe gtg	tat Tyr 180 aga	Asn 165 ttg Leu	643
96 97 99 100 101 103 104 105	Glu 150 act Thr gca Ala	gag Glu ttt	Phe att Ile caa Glr	Thr aga Arg gat Asp 185	gaa Glu 170 gtt Val	Met 155 gta Val ggt	Asp ggt Gly gct Ala	Leu cct Pro tgt	gtc Val gtt Val 190	Asp aac Asn 175 gcc	Arg 160 aag Lys ttg	Ile aag Lys gtg Val	Leu gga Gly g tct Ser	ttt Phe gtg Val	tat Tyr 180 aga	Asn 165 ttg Leu gta Val	643 691
96 97 99 100 101 103 104 105	Glu 150 act Thr gca Ala	gag Glu ttta Phe	Phe att Ile caa Glr aaa	Thr aga Arg agat Asp 185	gaa Glu 170 gtt Val	Met 155 gta Val ggt Gly	Asp ggt Gly gct Ala	cct Pro tgt Cys	gtc Val gtt Val 190 gtg	Asp aac Asn 175 gcc Ala	Arg 160 aag Lys ttg	aag Lys ggtg Val	gga gga g Gly g tct l Ser	ttt Phe tgtg Val 195	tat Tyr 180 aga Arg	Asn 165 ttg Leu gta Val	643
96 97 99 100 101 103 104 105 107	Glu 150 act Thr gca Ala	gag Glu ttta Phe	Phe att Ile Caa	Thr aga agata Arg Asp 185 aagg Lys	gaa Glu 170 gtt Val	Met 155 gta Val ggt Gly	Asp ggt Gly gct Ala	cct Pro tgt Cys aca	gtc Val gtt Val 190 gtg	Asp aac Asn 175 gcc Ala	Arg 160 aag Lys ttg	aag Lys ggtg Val	gga gga g Gly g tct l Ser g gct	ttt Phe y Phe y State y Val 195 y atg	tat Tyr 180 aga Arg	Asn 165 ttg Leu gta Val	643 691
96 97 99 100 101 103 104 105 107 108 109	Glu 150 act Thr gca Ala tac	gag Glu ttt Phe	Phe att Ile Caa Caa Caa Caa Caa Caa Caa Caa Caa Ca	Thr aga agat agat agat agat agag agag agag	gaa Glu 170 gtt Val	Met 155 gta Val ggt Gly	ggt Gly gct Ala ttt	cct Pro tgt Cys aca Thr	gtc Val gtt 190 gtg Val	aac Asn 175 gcc Ala aag	Arg 160 aag Lys Leu Leu aat	aag Lys gtg Val	gga gga g Gly g tct l Ser g gct n Ala 210	ttt Phe gtg Val 195 atg	tat Tyr 180 aga Arg	Asn 165 ttg Leu gta Val	643 691 739
96 97 99 100 101 103 104 105 107 108 109 111	Glu 150 act Thr gca Ala tac	gag Glu ttt Phe ttc Phe	Phe att Ile Case Glr aase Lys	Thr aga agat agat Asp 185 aaag s Lys	gaa Glu 170 gtt Val tgc Cys	Met 155 gta Val ggt Gly cca Pro	ggt Gly Gly Ala ttt	cct Pro Cys aca Thr 205	gtc Val gtt Val 190 gtg Val	Asp aac Asn 175 gcc Ala aag Lys	Arg 160 aag Lys Leu Asr	aag Lys gtg Val ctg Lev	gga gga g tct Ser gct Ala 210	ttt Phe y Phe y Val 195 y atg	tat Tyr 180 aga Arg	Asn 165 ttg Leu gta Val	643 691
96 97 99 100 101 103 104 105 107 108 109 111	Glu 150 act Thr gca Ala tac	gag Glu ttt Phe ttc Phe a cg	Phe att Ile Case Glr aas Lys 200	Thr aga agat agat Asp 185 aaag s Lys	gaa Glu 170 gtt Val tgc Cys	Met 155 gta Val ggt Gly cca Pro	ggt Gly Ala ttt	cct Pro Cys aca Thr 205	gtc Val gtt Val 190 gtg Val	Asp aac Asn 175 gcc Ala aag Lys	Arg 160 aag Lys Leu Asr	aag Lys ggtg Val ctg Leu gag	gga gga g tct Ser gct Ala 210 g gtt	ttt Phe y Phe y Val 195 y atg	tat Tyr 180 aga Arg	Asn 165 ttg Leu gta Val	643 691 739
96 97 99 100 101 103 104 105 107 108 109 111 112 113	Glu 150 act Thr gca Ala tac Tyr gac Asr	gag Gluatta Phe tto Phe acç Thi	Phe att I Caa Caa Caa Caa Caa Caa Caa Caa Caa C	Thr aga agata Asp 185 aaags Lys ccc	gaa Glu 170 gtt Val tgc Cys	Met 155 gta Val ggt Gly cca Pro	ggt Gly Ala ttt	cct Pro tgt Cys aca Thr 205	gtc Val gtt Val 190 gtg Val tcc Ser	aac Asn 175 gcc Ala aag Lys ctg	Arg 160 aag Lys Leu Leu Asr	aag Lys gete Val cete gege Glu	gga gga gtct Ser gct Ala 210 ggtt Val	ttt Phe Yal 195 Arg	tat Tyr 180 aga Arg	Asn 165 ttg Leu gta Val cca Pro	643 691 739 787
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115	Glu 150 act Thr gca Ala tac Tyr gac Asr	gag c Glu ttt a Phe c ttc c Phe c acc c Thi 21:	Phe att Ile Case Glr 200 gts Val	Thr aga agata Asp 185 aaag Lys Ccc Pro	gaa Glu 170 gtt Val tgc Cys atg	Met 155 gta Val ggt Gly cca Pro	ggt Gly Ala ttt Phe tcc Ser 220 gag	cct Pro tgt Cys aca Thr 205 cag	gtc Val gtt Yal 190 gtg Val tcc Ser	aac Asn 175 gcc Ala aag Lys ctg	Arg 160 aag Lys Leu Asr Val	aag Lys y gto val c cto i Lev gao Glu 225	gga gga gtct Ser gct Ala 210 gtt Val	ttt Phe Yal 195 Latg Arg	tat Tyr 180 aga Arg ttt Phe	Asn 165 ttg Leu gta Val cca Pro	643 691 739
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115 116	Glu 150 act Thr gca Ala tac Tyr gac Asr	gag c Glu ttt a Phe c ttc c Phe c acc c Thi 21:	Phe att Ile Case Glr 200 gts Val	Thr aga agata Asp 185 aaag Lys Ccc Pro	gaa Glu 170 gtt Val tgc Cys atg	Met 155 gta Val ggt Gly cca Pro	ggt Gly Ala ttt Phe tcc Ser 220 gag	cct Pro tgt Cys aca Thr 205 cag	gtc Val gtt Yal 190 gtg Val tcc Ser	aac Asn 175 gcc Ala aag Lys ctg	Arg 160 aag Lys Leu Asr Val	aag Lys y gto val c cto i Lev gao Glu 225	gga gga gtct Ser gct Ala 210 gtt Val	ttt Phe Yal 195 Latg Arg	tat Tyr 180 aga Arg ttt Phe	Asn 165 ttg Leu gta Val cca Pro	643 691 739 787
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115	Glu 150 act Thr gca Ala tac Tyr gac Asr	gag Glu ttt Phe c ttc Phe c acc Thi 215 c gtc Val	Phe att Ile Case Glr aase Lys 200 gts Val	Thr aga agata Asp 185 aaag Lys Ccc Pro	gaa Glu 170 gtt Val tgc Cys atg	Met 155 gta Val ggt Gly cca Pro	ggt Gly Gly Ala ttt Phe tcc Ser 220 gag	cct Pro tgt Cys aca Thr 205 cag	gtc Val gtt Yal 190 gtg Val tcc Ser	aac Asn 175 gcc Ala aag Lys ctg	Arg 160 aag Lys Leu Asr Val	aag s Lys y gtg val c ctg s Leu g gag c Glu 225	gga gga gtct Ser gct Ala 210 gtt Val	ttt Phe Yal 195 Latg Arg	tat Tyr 180 aga Arg ttt Phe	Asn 165 ttg Leu gta Val cca Pro	643 691 739 787
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115 116	Glu 150 act Thr gca Ala tac Tyr gac Asr	gag c Glu ttt a Phe c ttc c Phe c acc c Thi 215 c gtc s Val	The att of the control of the contro	Thr aga aga agat agat agat agag agag agag	gaa Glu 170 gtt Val tgc Cys atg	Met 155 gta Val ggt Gly cca Pro gac Asp	ggt Gly Ala ttt Phe tcc Ser 220 gag	cct Pro tgt Cys aca Thr 205 cag Gln	gtc Val gtt Val 190 gtg Val tcc Ser gat Asp	aac Asn 175 gcc Ala aag Lys ctg	Arg 160 aag Lys Leu Asr Val	aag s Lys y gtg val c ctg s Leu g gag c Glu 225	gga gga gtct Ser gct Ala 210 gtt Val g atg	ttt Phe y Phe y Val 195 atg Met Arg	tat Tyr 180 aga Arg ttt Phe Gly	Asn 165 ttg Leu gta Val cca Pro tct Ser agt Ser 245	643 691 739 787
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115 116 117	Glu 150 act Thr gca Ala tac Tyr gac Asr	gag c Glu ttt a Phe c ttc c Phe c acc c Th 215 c gtc c Val	The att of the control of the contro	Thr aga agat agat agat agag agag agag agag	gaa Glu 170 gtt Val tgc Cys atg	Met 155 gta Val ggt Cca Pro gac Asp Lys 235	ggt Gly gct Ala ttt Phe tcc Ser gag	cct Pro tgt Cys aca Thr 205 cag Gln	gtc Val gtt Val 190 gtg Val tcc Ser gat Asp	aac Asn 175 gcc Ala aag Lys ctg	Arg 160 aag Lys Leu Asr yal Pro 240 aag	aag S Lys S Lys S Lys S S Lys S S S S S S S S S S S S S S S S S S S	gga gga gtct Ser gct Ala 210 gtt Val gtt Met	ttt Phe gtg Val 195 atg Met Arg	tat Tyr 180 aga Arg ttt Phe Gly Cys	Asn 165 ttg Leu gta Val cca Pro tct Ser 245 gct	643 691 739 787 835
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115 116 117 119	Glu 150 act Thr gca Ala tac Tyr gac Asr	gag c Glu ttt a Phe c ttc c Phe c acc c Th 215 c gtc c Val	The att of the control of the contro	Thr aga agat agat agat agag agag agag agag	gaa Glu 170 gtt Val tgc Cys atg	Met 155 gta Val ggt Gly cca Pro gac Asp Lys 235 ctt	ggt Gly gct Ala ttt Phe tcc Ser gag	cct Pro tgt Cys aca Thr 205 cag Gln	gtc Val gtt Val 190 gtg Val tcc Ser gat Asp	aac Asn 175 gcc Ala aag Lys ctg	Arg 160 aag Lys Leu Asn yal Pro 240 aag Lys	aag S Lys S Lys S Lys S S Lys S S S S S S S S S S S S S S S S S S S	gga gga gtct Ser gct Ala 210 gtt Val gtt Met	ttt Phe gtg Val 195 atg Met Arg	tat Tyr 180 aga Arg ttt Phe Gly Cys aat	Asn 165 ttg Leu gta Val cca Pro tct Ser 245 gct Ala	643 691 739 787 835
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115 116 117 119 120 121	Glu 150 act Thr gca Ala tac Tyr gac Asr tgt Cys 230 aca Thr	gag Glu ttt a Phe c ttc c Phe c acc c Th 215 c gtc c Val d gaa c Glu	Phe att I I I I I I I I I I I I I I I I I I	aga agat agat agat agat agas agag agag a	gaa Glu 170 gtt Val Cys atg Met tct Ser tgg Trp 250	Met 155 gta Val ggt Gly cca Pro Asp Lys 235 ctt Leu	ggt Gly Ala ttt Phe tcc Ser 220 gag Glu ttt Val	cct Pro tgt Cys aca Thr 205 cag Gln gaa Glu	gtc Val gtt Val 190 gtg Val tcc Ser gat Asp	Asp aac Asn 175 gcc Ala agg Lys ctg Cet Gly 255	Arg 160 aag Lys Leu Asn Yal Cca Pro 240 aag Lys	aag s Lys s Lys s Cys s Cys	gga gga g tct Ser gct Ala 210 gtt val g atg Met	ttt Phe gtg Val 195 Arg Tyr ctgc	tat Tyr 180 aga Arg ttt Phe Gly Cys aat Asn	Asn 165 ttg Leu gta Val cca Pro tct Ser 245 gct Ala	643 691 739 787 835
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115 116 117 119 120 121 123	Glu 150 act Thr gca Ala tac Tyr gac Asp tgt Cys 230 aca Thr	gag Glu ttt a Phe c ttc C Th 215 C Yal a gaa c Glu c tat	Phe att I le Caa e Glr 200 gta Caa caa ca caa ca caa ca caa ca caa ca	Thr aga agat agat agat agat agag agag agaa agaa	gaa 170 gtt Val tgc Cys Atg Met tct Ser tgg Trp 250 aga	Met 155 gta Val ggt Gly cca Pro Asp Lys ctt Leu ggt	ggt gct Ala ttt Phe cc Ser 220 gag Glu cc gtal Val	cct Pro tgt Cys aca Thr 205 cag Gln gaa Glu ccc	gtc Val gtt Val 190 gtg Val tcc Ser gat Asp	aac Asn 175 gcc Ala aag Lys ctg Leu cct ggc	Arg 160 aag Lys Leu Asn yal Cca Pro 240 aag Lys	aag s Lys g gtg val c ctg s Lev g gag s Arg t tgt	gga gga g tct Ser gct Ala 210 gtt Val g atg	ttt Phe y Phe y Val 195 atg Arg Tyr tgc Cys	tat Tyr 180 aga Arg ttt Phe Gly Cys aat Asr	Asn 165 ttg Leu gta Val cca Pro tct Ser 245 gct Ala	643 691 739 787 835
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115 116 117 119 120 121 123 124	Glu 150 act Thr gca Ala tac Tyr gac Asp tgt Cys 230 aca Thr	gag Glu ttt a Phe c ttc C Th 215 C Yal a gaa c Glu c tat	Phe att I le Caa e Glr 200 gta Caa caa ca caa ca caa ca caa ca caa ca	Thr aga gat Asp 185 aags Lys Coc Pro aat Asn Glu gaa Glu	gaa 170 gtt Val tgc Cys Atg Trp 250 aga Arg	Met 155 gta Val ggt Gly cca Pro Asp Lys ctt Leu ggt	ggt gct Ala ttt Phe cc Ser 220 gag Glu cc gtal Val	cct Pro tgt Cys aca Thr 205 cag Gln gaa Glu ccc	gtc Val gtt Val 190 gtg Val tcc Ser gat Asp	aac Asn 175 gcc Ala aag Lys ctg Cot Gly 255 caa Gln	Arg 160 aag Lys Leu Asn yal Cca Pro 240 aag Lys	aag s Lys g gtg val c ctg s Lev g gag s Arg t tgt	gga gga g tct Ser gct Ala 210 gtt Val g atg	ttt Phe gtg Val 195 Arg Tyr Cys	tat 180 aga Arg ttt Phe Gly Cys aat Asn 260 ggt	Asn 165 ttg Leu gta Val cca Pro tct Ser 245 gct Ala	643 691 739 787 835
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115 116 117 119 120 121 123 124 125	Glu 150 act Thr gca Ala tac Tyr gac Asr tgt Cys 230 aca Thr	gag Glu ttt a Phe c tto C Phe c acc C Thi 215 c gto C Glu c tat Tyi	Phe att I le Caa le Glr Vall Asr Glr Glr Glr	Thr aga gat Asp 185 aag CCC Pro aat Asn Glu gaa Glu 265	gaa 170 gtt Val tgc Cys Atg Trp 250 aga Arg	Met 155 gta Val ggt Gly Cca Pro Asp Lys 235 ctt Leu ggt	ggt Gly Ala ttt Phe	cct Pro tgt Cys aca Thr 205 cag Gln gaa Glu ccc Pro	gtc Val gtt Val 190 gtg Val tcc Ser gat att	aac Asn 175 gcc Ala aag Lys ctg Leu cct gcc Gly 255 caa Gln	Arg 160 aag Lys Leu Asr Val Cca Pro 240 aag Lys Ala	aag S Lys S Lys S Cys S Cys C Cys	gga gga ggt ggt l Ser ggt l Ala 210 ggt l Val g atg g Arg	ttt Phe y Phe y Val 195 Latg Met Arg tac Tyr Cys	tat 180 aga Arg ttt Phe Gly cys aat Ass 260 ggt	Asn 165 ttg Leu gta Val cca Pro tct Ser 245 gct Ala	643 691 739 787 835 883
96 97 99 100 101 103 104 105 107 108 109 111 112 113 115 116 117 119 120 121 123 124	Glu 150 act Thr gca Ala tac Tyr gac Asr tgt Cys 230 aca Thr	gag Glu ttta Phe c ttc Phe c acc So Thi 215 C gtc So Val n gaa c Glu c tat y Typ	Phe att I I I I I I I I I I I I I I I I I I	Thr aga gat Asp 185 aaag CCC Asn Glu gaa Glu 265 ttg	gaa Glu 170 gtt Val tgc Cys Atg Trp 250 aga Arg	Met 155 gta Val ggt Gly cca Pro Asp Lys 235 ctt Gly ggt	ggt Gly Ala ttt Phe Con Grand Glu Tttt Phe Con Grand Glu Tttt Ttt Phe Con Grand Glu Tttt The Grand Glu Tttt Ttt The Grand Glu Tttt Ttt Ttt Ttt Ttt Ttt Ttt Ttt Ttt T	cct Pro tgt Cys aca Thr 205 cag Gln gaa Glu ccc Pro	gtc Val gtt Val 190 gtg Val tcc Ser gat Asp atte Cys aag	aac Asp 175 gcc Ala aag Lys ctg Cot Gly 255 caaa Gln tgt	Arg 160 aag Lys Leu Asr Val Pro 240 aag Lys Asr Lys Gaag	aag Lys y gtg Val Let gag Arg Arg y tgt	gga gga gtct Ser gct Ala 210 gtt Val gtt Ser cca Arg	ttt Phe Yell Yell Yell Yell Yell Yell Yell Ye	tat Tyr 180 aga Arg ttt Phe Gly tgggt Gly Gly Gly cct	Asn 165 ttg Leu gta Val cca Pro tct Ser 245 gct Ala	643 691 739 787 835

Input Set : A:\PTO.AMC.txt

129			280					285					290				
131	24+	tat		a = a	~~~	an t	aat		atg	220	+~~	200		a a a	2 2 t	22t	1027
																	1027
132	Ser		TIIT	GIII	GIU	ASP		Ser	Met	ASII	Cys		Cys	GIU	ASII	ASII	
133	.	295	~~~	~~~	~~~		300	+		+	~+~	305	+~+		-~-		1075
135				_	_		_		cca		_	-	_		_		1075
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139					_		_		tct							-	1123
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141					330					335					340		
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144	Ile	Leu	Asp	Trp	Ser	Trp	Pro	Leu	Asp	Thr	Gly	Gly	Arg	Lys	Asp	Val	
145				345					350					355			
147									tgt								1219
148	Thr	Phe	Asn	Ile	Ile	Cys	Lys	Lys	Cys	Gly	${\tt Trp}$	Asn	Ile	Lys	Gln	Cys	
149			360					365					370				
151	gag	cca	tgc	agc	cca	aat	gtc	cgc	ttc	ctc	cct	cga	cag	ttt	gga	ctc	1267
152	Glu	Pro	Cys	Ser	Pro	Asn	Val	Arg	Phe	Leu	Pro	Arg	Gln	Phe	Gly	Leu	
153		375					380					385					
155	acc	aac	acc	acg	gtg	aca	gtg	aca	gac	ctt	ctg	gca	cat	act	aac	tac	1315
156	Thr	Asn	Thr	Thr	Val	Thr	Val	Thr	Asp	Leu	Leu	Ala	His	Thr	Asn	Tyr	
157	390					395					400					405	
159	acc	ttt	gag	att	gat	qcc	gtt	aat	ggg	gtg	tca	gag	ctg	agc	tcc	cca	1363
160									Gly								
161					410				•	415					420		
163	cca	aga	cag	ttt	act	aca	atc	agc	atc	aca	act	aat	caq	act	act	cca	1411
164		-	_		-		_	_	Ile				_	-	_		
165		5	0	425					430					435			
167	t.ca	cct.	at.c		acσ	att.	aaσ	aaa	gat	caa	acc	t.cc	aσa	aat.	aσc	atc	1459
168			-	_	_		_		-				_		-	Ile	
169			440				-1-	445		5			450				
171	tet	tta		t.aa	caa	αаа	aat.		cat	cct.	aat	aaa		ata	t.t.a	gac	1507
172		_				_		_	His						_	-	
173	001	455	501		01	014	460	014				465					
175	tac		atc	aaa	tac	tat		ааσ	cag	αаа	caa		aca	aαt	tat	acc	1555
176									Gln								1333
177	470	Olu	, 41		-1-	475	OLU		0111	014	480	Olu		001	-1-	485	
179		ata	200	aca	2012		202	aat	gtt	200		a art	200	ata	224		1603
180									Val								1003
181	116	пеп	лту	AIG	490	GLY	1111	ASII	Val	495	116	Set	SET	пеп	500	PIO	
183	~~~	20+	2+2	+ 2.0		++0	633	a+c	002		003	202	~~~	act		+ > +	1651
184									cga Arg								1031
185	ASP	TIIT	116	505	vaı	Pne	GIII	116	510	мта	Arg	1111	нта		СТУ	TAT	
														515			1600
187									ttt								1699
188	ĠΤĀ	Tnr		ser	Arg	тĀ2	rne		Phe	GIU	rnr	ser		Asp	ser	ьие	
189	.		520					525		4	_4 -		530				
191									gtg								1747
192	ser		ser	GTĀ	GLu	ser		GIn	Val	val	мet		ΑΙα	тте	ser	Ата	
193		535					540					545					

Input Set : A:\PTO.AMC.txt

																1795
	Val	Ala	Ile	Ile		Leu	Thr	Val	Val		Tyr	Val	Leu	Ile	-	
	++-					+										1042
																1843
Arg	Pne	Cys	СТА	_	гуѕ	ser	гаг	HIS	-	Ата	Asp	GIU	гля	_	Leu	
ast	+++	aac	22+		cat	++2	222	att		aat	ata	200	20+		a++	1001
															-	1891
птэ	FIIC	СТУ		GTĀ	птэ	пец	цуз		PIO	СТУ	Бец	Aly		тут	vai	
gac	cca	cat		tat	gaa	gac	cct		саа	act	att	cat		+++	acc	1939
																1757
		600		-1-			605		0111		, 42		014	1110	1114	
aag	gaa	ttg	gat	gcc	acc	aac	ata	tcc	att	gat	aaa	qtt	qtt	qqa	qca	1987
_	-	_	-	-						_		-	-		-	
-	615		_			620				_	625			-		
ggt	gaa	ttt	gga	gag	gtg	tgc	agt	ggt	cgc	tta	aaa	ctt	cct	tca	aaa	2035
Gly	Glu	Phe	Gly	Glu	Val	Cys	Ser	Gly	Arg	Leu	Lys	Leu	Pro	Ser	Lys	
630					635					640					645	
																2083
Lys	Glu	Ile	Ser		Ala	Ile	Lys	Thr		Lys	Val	Gly	\mathtt{Tyr}		Glu	
									655							
																2131
Lys	Gln	Arg	_	Asp	Phe	Leu	Gly		Ala	Ser	Ile	Met	_	Gln	Phe	
																0170
_						_	_	_		_	-			_	_	2179
Asp	HIS		ASI	тте	ше	Arg		GIU	GIA	Val	vaı		гаг	ser	гÀг	
003	~++		2++	a+a	202	ma a		2+4	a a a	a a +	aat		++~	ant.	244	2227
																2221
FIO		Mec	116	val	1111		TYT	Mec	GIU	HOII		361	пеп	nsp	261	
ttc		cat	aaa	cac	gat		cad	+++	act	atc		саσ	cta	ata	aaa	2275
		_			-	_	_			_		_				2273
710		5	-1-		715					720					725	
atq	ctt	cqa	qqq	ata	gca	tct	qqc	atq	aaq	tac	ctq	tca	qac	atq	qqc	2323
_					_			_	_		_		_	_		
				730					735	_				740	_	
tat	gtt	cac	cga	gac	ctc	gct	gct	cgg	aac	atc	ttg	atc	aac	agt	aac	2371
Tyr	Val	His	Arg	Asp	Leu	Ala	Ala	Arg	Asn	Ile	Leu	Ile	Asn	Ser	Asn	
			745					750					755			
																2419
Leu	Val	_	Lys	Val	Ser	Asp		Gly	Leu	Ser	Arg		Leu	Glu	Asp	
																2467
Asp		Glu	Ala	Ala	Tyr		Thr	Arg	Gly	Gly	-	Ile	Pro	Ile	Arg	
•																
																2515
_	rnr	ser	PLO	GIU		тте	ата	Tyr	arg	_	ьиe	rnr	ser	Ата		
	at s	+~~	a ~+	+-+		a++	~++	at a	+~~		a+~	a+~	+ <+	+=+		2562
yaı	yıa	Lyy	ayı	Lat	999	all	gıı	CLC	Lyg	gag	gicg	alg	LUL	Lat	yya	2563
	Ala 550 agg cat his gac Asp agg tys aag Lys gaty gac Asp cao ttephe 710 atg tteu gac Asp tgp 790	Ala Val 550 agg ttc Arg Phe cat ttt His Phe gac Cca Asp Pro aag gaa Lys Glu 630 aaa gag Lys Glu aag Cag Lys Glu aag Cag Lys Glu aag Cag Lys Glu atg Cac Asp His cca gtt Pro 695 ttc Cta Phe Cta Phe Leu 710 atg tt Met Leu tat gtt Tyr Val ttg gtg Leu caa Asp Pro ttg gtg Thr 790	Ala Val Ala 550 agg ttc tgt Arg Phe Cys cat ttt ggc His Phe Gly gac cca cat Asp Pro His 600 aag gaa ttg Lys Glu Leu 615 ggt gaa ttt Gly Glu Phe 630 aaa gag att Lys Glu Ile aag cag agg Lys Gln Arg gac cac ccc Asp His Pro 680 cca gtt atg Pro Val Met 695 ttc cta cgt Phe Leu Arg 710 atg ctt cga Met Leu Arg 710 atg ctt cga Met Leu Arg 710 atg ctt cac Tyr Val His ttg gtg tgt Leu Val Cys 760 gac cca gaa Asp Pro tgg aca tca Trp Thr 790	Ala Val Ala Ile 550 agg ttc tgt ggc Arg Phe Cys Gly cat ttt ggc aat His Phe Gly Asn 585 gac cca cat aca Asp Pro His Thr 600 aag gaa ttg gat Lys Glu Leu Asp 615 ggt gaa ttt gga Gly Glu Phe Gly 630 aaa gag att tca Lys Glu Ile Ser aag Cag agg aga Lys Gln Arg Arg 665 gac cac act Asp His Pro Asn 680 cca gtt atg Arg 665 gac cac ccc aat Asp His Pro Asn 680 cca gtt atg att Pro Val Arg 680 cta gtt atg att Pro Asn 680 cta aaa Pro Asn 680 cta aaa Arg Arg 665 aaa Arg	Ala Val Ala Ile Ile 550 ttt tgg tat agg ttc tgg ggc tat Arg Phe Cys Gly Tyr 570 cat ttt ggc aat ggg His Phe Gly Asn Gly 585 gac cat aca tat Asp Pro His Thr Tyr 600 aag gcc Lys Ala 615 Thr Tyr Asp Ala 615 Leu Asp Ala 630 Thr Tyr Asp 631 Thr Tyr Asp 632 Gau ttc gg gag 633 att tca gtg 645 agg aga gac Lys Glu Arg Asp 650 aag aga gac Asp Att gc gc Asp<	Ala Ile Leu 550 to to 555 agg ttc tgt ggc tat aag Arg Phe Cys Gly Tyr Lys cat ttt ggc aat gga cat gac cca cat aca tga ga gac cca cat aca tat ga gac cca cat aca tat ga gac cca tat dac acc Lys Glu Leu Asp Ala Thr gdt gaa ttt gga ggg gtg gdt gaa ttt cac ggg gtg gdt gaa ttt cac ggg ttt gdt atg aca tt tt gdt atg aca tt gac cac aca	Ala Val Ala Ile Leu Leu 555 agg ttc tgt ggc tat aag tca Arg Phe Cys Gly Tyr Lys Ser 570 Tyr Lys Ser 570 Tyr Lys Ser 6at ttt Asp Gly His Leu Ser 570 Tyr Lys Ser gac cca Cat Asp Gly His Leu Asp His Leu Asp His Leu Asp Asp Glu Asp Asp	Ala Val Lee Lee Thr 550 The Cot Gly Tyr Lys Seer Lys agg ttt ggc aat aag tca aaa Arg Phe Cys Gly Tyr Lys Seer Lys cat ttt ggc cat tag aaa aaa gac cca cat ggy cat tag gac cct fasp Pro His Thr Tyr Glu Asp Pro 605 aag gaa ttg gat gac act aaa <	Ala Val Ile Leu Leu Thr Val 555 tca taag tca aaa cat Arg Phe Cys Gly Tyr Lys Ser Lys His cat ttt ggg cat tta aaa ctt His Phe Gly Asn Gly His Leu Lys Leu Asp Pro His Thr Tyr Glu Asp Pro Thr 605 aag gaa ttg gat gcc acc aca aca acc Asp Pro Thr 605 Thr 610 Thr 610 Thr 610 Thr 610 Thr 610 Thr 610 <	Ala Val Ala Ile Leu Leu Thr Val Val 550 Ty Sep tat aag tca aaa cat ggg Arg Phe Cys Gly Tyr Lys Ser Lys His Gly 575 cat ttt ggg cat tta aaa ctt cca dat cot 575 cot 575 575 cot 575 575 575 cot 575	Ala Val Ala Ile Leu Leu Thr Val Val Ile 550 555 560 560 560 560 360 360 Arg Phe Cys Gly Tyr Lys Eu Lys His Gly Ala 570 cat ttt gg cat ttt aaa ctt cac ggt pro FTD FTD	Ala Val Ala Ile Ile Leu Leu Thr Val Val Ile Tyr 550 555 555 555 560	Ala of the state of t	Ala Val Ala Ile Ile Leu Leu Thr Val Val Ile Tyr Val Leu Sep Sep	Ala Val Ala Ile Ile Leu Leu Thr Val Val Ile Tyr Val Ile Ile S55 S55	Arg Phe Cys Gly Tyr Lys Ser Lys His Gly Ala Asp Glu Lys Arg Leu Ser Sys Sys Ser Lys His Gly Ala Asp Glu Lys Arg Leu Sys

Input Set : A:\PTO.AMC.txt

0.00		**- 7				01	-1-		-		a 1.		34.4	a	_	a 1.		
260	Asp	Val	тгр	ser	810	Gly	TTE	vaı	Leu	815	GIU	vaı	met	ser	820	GIY		
261 263	a a a	242	002	tac		gag	a t or	+00	22t		ra+	at a	2++	222		at a		2611
264		-				Glu	_			_	_	_			-	-		2011
265	Giu	AIG	FIU	825	115	Giu	nec	DCI	830	GIII	nsp	Val	116	835	AIG	Val		
267	gat	σασ	aac	-	cga	ctg	cca	CCC		atσ	gac	tac	сса		acc	tta	•	2659
268	_				-	Leu				_	-	_		-	-	_		2000
269	PF	52 4	840	-1-	9			845				0,0	850					
271	tat	caq	cta	atq	cta	gac	tac		caq	aaa	gac	agg	aac	aac	aσa	ccc		2707
272		_	_	_	_	Asp	_		_		_				_			
273	-	855				•	860	-		-	•	865			•			
275	aag	ttt	gag	cag	att	gtt	agt	att	ctg	gac	aag	ctt	atc	cqq	aat	ccc		2755
276						Val												
277	870					875				-	880					885		
279	ggc	agc	ctg	aag	atc	atc	acc	agt	gca	gcc	gca	agg	cca	tca	aac	ctt		2803
280	Gly	Ser	Leu	Lys	Ile	Ile	Thr	Ser	Ala	Āla	Ala	Arg	Pro	Ser	Asn	Leu		
281					890					895					900			
283	ctt	ctg	gac	caa	agc	aat	gtg	gat	atc	tct	acc	ttc	cgc	aca	aca	ggt		2851
284	Leu	Leu	Asp	Gln	Ser	Asn	Val	Asp	Ile	Ser	Thr	Phe	Arg	Thr	Thr	Gly		
285				905					910					915				
287						gtc												2899
288	Asp	\mathtt{Trp}	Leu	Asn	Gly	Val	Arg	Thr	Ala	His	Cys	Lys	Glu	Ile	Phe	Thr		
289			920					925					930					
291					_	tct	_	_			_	_				_		2947
292	Gly		Glu	Tyr	Ser	Ser	_	Asp	Thr	Ile	Ala	_	Ile	Ser	Thr	Asp		
293		935					940					945						
295						ggt												2995
296	_	Met	Lys	Lys	Val	Gly	Val	Thr	Val	Val	_	Pro	Gln	Lys	Lys			
297	950					955					960					965		
299						gct												3043
300	шe	Ser	ser	тте		Ala	Leu	Glu	Thr		ser	Lys	Asn	GTĀ		Val		
301					970					975					980			2002
303 304	Pro		taa	agca	acga	cgg a	aagt	gette	et ge	gacg	gaagi	c ggi	Eggei	_g c g				3092
304				+	.+	٠~ ~.		. ~ ~ ~ .	+.	++	a+ ~ ~	2021	- a		~~~~			2140
				_		tg ca	agaca	iyaca	aac	aatt	cugg	ayaı	Lacti	gg L s	yyaaq	JLL		3149
	<210×																	
	<212				,													
	<213				Omo	ean.	iane											
	<400					bup.	LCIIS											
315						Ser	Tle	T.e.11	Leu	T.e.11	T.e.11	Ser	Cvs	Ser	Va 1	T.A11		
316	1		Cyb	0111	5	DCI	110	Dea	Lea	10	пси	UCI	CIS	DCI	15	Deu		
317	_	Ser	Phe	Glv		Leu	Ile	Pro	Gln		Ser	Asn	Glu	Va 1		Len		
318				20					25					30				
319	Leu	Asp	Ser		Thr	Ile	Gln	Glv		Leu	Glv	Trp	Ile		Tvr	Pro		
320			35	_1 ~				40			1		45		-1-			
321	Ser	His		Trp	Glu	Glu	Ile		Glv	Val	Asp	Glu		Tyr	Thr	Pro		
322		50	- 4				55		- 4		- 1	60			J	-		
323	Ile	Arg	Thr	Tyr	Gln	Val	Cys	Asn	Val	Met	Asp	His	Ser	Gln	Asn	Asn		
		-		-			-				_							

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/913,756

DATE: 03/07/2002 TIME: 18:40:44

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF3\03072002\I913756.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the $\langle 220 \rangle$ to $\langle 223 \rangle$ fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 43